SEQUENCE LISTING <110> Christopher D Castle, John C Garrett-Engele, Philip W Kan, Zhengyan Loerch, Patrick M Tsinoremas, Nicholas F <120> Novel Isoforms of Centromere Protein E (CENPE) <130> RS0210Y <140> US 10/828,985 <141> 2004-04-21 <150> US 60/464,905 <151> 2003-04-23 <150> US 60/510,701 <151> 2003-10-10 <160> 25 <170> PatentIn version 3.2 <210> 1 <211> 40 <212> DNA <213> Homo sapiens <400> 1 aagatgaatt acagaaaaag atccaagaac ttcagaaaaa 40 <210> 2 <211> 40 <212> DNA <213> Homo sapiens <400> 2 taagggaaat gatagctaga gaccgacaga accaccaagt 40 <210> 3 <211> 40 <212> DNA <213> Homo sapiens <400> 3 aagatgaatt acagaaaaag gaccgacaga accaccaagt 40 <210>

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Thr Ile Phe Ala Tyr Gly Gln Thr Ala Ser Gly Lys Thr Tyr Thr Met 85 90 95

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Val Ser Tyr Met Glu Ile Tyr Asn Glu Thr Ile Thr Asp Leu Leu Cys 130 135 140

Gly Thr Gln Lys Met Lys Pro Leu Ile Ile Arg Glu Asp Val Asn Arg 145 150 155 160

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Pro	Ser 2090	Arg	Glu	Leu	Arg	Asp 2095	Leu	Lys	Leu	Asn	Gln 2100	Asn	Met	Asp
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Leu	Phe 2210	Lys	Asn	Tyr	Gln	Thr 2215	Leu	Lys	Thr	Ser	Leu 2220	Ala	Ser	Gly
Ala	Gln 2225	Val	Asn	Pro	Thr	Thr 2230	Gln	Asp	Asn	Lys	Asn 2235	Pro	His	Val
Thr	Ser 2240	Arg	Ala	Thr	Gln	Leu 2245	Thr	Thr	Glu	Lys	Ile 2250	Arg	Glu	Leu
Glu	Asn 2255	Ser	Leu	His	Glu	Ala 2260	Lys	Glu	Ser	Ala	Met 2265	His	Lys	Glu
Ser	Lys 2270	Ile	Ile	Lys	Met	Gln 2275	Lys	Glu	Leu	Glu	Val 2280	Thr	Asn	Asp
Ile	Ile 2285	Ala	Lys	Leu	Gln	Ala 2290	Lys	Val	His	Glu	Ser 2295	Asn	Lys	Cys
Leu	Glu 2300	Lys	Thr	Lys	Glu	Thr 2305	Ile	Gln	Val	Leu	Gln 2310	Asp	Lys	Val
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Leu Gly Leu Ser Lys Arg Leu Gln Glu Ser His Asp Glu Met Lys Ser Val Ala Lys Glu Lys Asp Asp Leu Gln Arg Leu Gln Glu Val Leu Gln Ser Glu Ser Asp Gln Leu Lys Glu Asn Ile Lys Glu Ile Val Ala Lys His Leu Glu Thr Glu Glu Glu Leu Lys Val Ala His Cys Cys Leu Lys Glu Gln Glu Glu Thr Ile Asn Glu Leu Arg Val Asn Leu Ser Glu Lys Glu Thr Glu Ile Ser Thr Ile Gln Lys Gln Leu Glu Ala Ile Asn Asp Lys Leu Gln Asn Lys Ile Gln Glu Ile Tyr Glu Lys Glu Glu Gln Leu Asn Ile Lys Gln Ile Ser Glu Val Gln Glu Asn Val Asn Glu Leu Lys Gln Phe Lys Glu His Arg Lys Ala Lys Asp Ser Ala Leu Gln Ser Ile Glu Ser Lys Met Leu Glu 1535 1540 Leu Thr Asn Arg Leu Gln Glu Ser Gln Glu Glu Ile Gln Ile Met 1550 1555 Ile Lys Glu Lys Glu Glu Met Lys Arg Val Gln Glu Ala Leu Gln Ile Glu Arg Asp Gln Leu Lys Glu Asn Thr Lys Glu Ile Val Ala Lys Met Lys Glu Ser Gln Glu Lys Glu Tyr Gln Phe Leu Lys Met

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Lys Lys Val Ser Glu Met Glu Gln Leu Lys Lys Gln Ile Lys Asp Gln Ser Leu Thr Leu Ser Lys Leu Glu Ile Glu Asn Leu Asn Leu Ala Gln Glu Leu His Glu Asn Leu Glu Glu Met Lys Ser Val Met Lys Glu Arg Asp Asn Leu Arg Arg Val Glu Glu Thr Leu Lys Leu Glu Arg Asp Gln Leu Lys Glu Ser Leu Gln Glu Thr Lys Ala Arg Asp Leu Glu Ile Gln Gln Glu Leu Lys Thr Ala Arg Met Leu Ser Lys Glu His Lys Glu Thr Val Asp Lys Leu Arg Glu Lys Ile Ser Glu Lys Thr Ile Gln Ile Ser Asp Ile Gln Lys Asp Leu Asp Lys Ser Lys Asp Glu Leu Gln Lys Lys Asp Arg Gln Asn His Gln Val Lys Pro Glu Lys Arg Leu Leu Ser Asp Gly Gln Gln His Leu Met Glu Ser Leu Arg Glu Lys Cys Ser Arg Ile Lys Glu Leu Leu Lys Arg Tyr Ser Glu Met Asp Asp His Tyr Glu Cys Leu Asn Arg Leu Ser Leu Asp Leu Glu Lys Glu Ile Glu Phe His Arg Ile Met Lys Lys Leu Lys Tyr Val Leu Ser Tyr Val Thr Lys Ile Lys Glu Glu

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Ala Met His Lys Glu Ser Lys Ile Ile Lys Met Gln Lys Glu Leu Glu Val Thr Asn Asp Ile Ile Ala Lys Leu Gln Ala Lys Val His Glu Ser Asn Lys Cys Leu Glu Lys Thr Lys Glu Thr Ile Gln Val Leu Gln Asp Lys Val Ala Leu Gly Ala Lys Pro Tyr Lys Glu Glu . Ile Glu Asp Leu Lys Met Lys Leu Val Lys Ile Asp Leu Glu Lys 2300 2305 Met Lys Asn Ala Lys Glu Phe Glu Lys Glu Ile Ser Ala Thr Lys Ala Thr Val Glu Tyr Gln Lys Glu Val Ile Arg Leu Leu Arg Glu Asn Leu Arg Arg Ser Gln Gln Ala Gln Asp Thr Ser Val Ile Ser Glu His Thr Asp Pro Gln Pro Ser Asn Lys Pro Leu Thr Cys Gly Gly Gly Ser Gly Ile Val Gln Asn Thr Lys Ala Leu Ile Leu Lys Ser Glu His Ile Arg Leu Glu Lys Glu Ile Ser Lys Leu Lys Gln Gln Asn Glu Gln Leu Ile Lys Gln Lys Asn Glu Leu Leu Ser Asn Asn Gln His Leu Ser Asn Glu Val Lys Thr Trp Lys Glu Arg Thr Leu Lys Arg Glu Ala His Lys Gln Val Thr Cys Glu Asn Ser Pro

Lys Ser Pro Lys Val Thr Gly Thr Ala Ser Lys Lys Gln Ile 2450 2455 2460 Thr Pro Ser Gln Cys Lys Glu Arg Asn Leu Gln Asp Pro Val Pro 2465 2470 2475 Lys Glu Ser Pro Lys Ser Cys Phe Phe Asp Ser Arg Ser Lys Ser 2480 2485 2490 Leu Pro Ser Pro His Pro Val Arg Tyr Phe Asp Asn Ser Ser Leu 2495 2500 2505 Gly Leu Cys Pro Glu Val Gln Asn Ala Gly Ala Glu Ser Val Asp 2510 Ser Gln Pro Gly Pro Trp His Ala Ser Ser Gly Lys Asp Val Pro 2525 2530 Glu Cys Lys Thr Gln 2540 <210> 10 <211> 7509 <212> DNA <213> Homo sapiens <400> 10 atggcggagg aaggagccgt ggccgtctgc gtgcgagtgc ggccgctgaa cagcagagaa 60 gaatcacttg gagaaactgc ccaagtttac tggaaaactg acaataatgt catttatcaa 120 gttgatggaa gtaaatcctt caattttgat cgtqtctttc atgqtaatqa aactaccaaa 180 aatgtgtatg aagaaatagc agcaccaatc atcgattctg ccatacaagg ctacaatggt 240 actatatttg cctatggaca gactgcttca ggaaaaacat ataccatgat gggttcagaa 300 gatcatttgg gagttatacc cagggcaatt catgacattt tccaaaaaat taaqaaqttt 360 cctgataggg aatttctctt acgtgtatct tacatggaaa tatacaatga aaccattaca 420 gatttactct gtggcactca aaaaatgaaa cctttaatta ttcqaqaaqa tqtcaataqq 480 aatgtgtatg ttgctgatct cacagaagaa gttgtatata catcagaaat qqctttgaaa 540

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Met Gly Ser Glu Asp His Leu Gly Val Ile Pro Arg Ala Ile His Asp 100 105 110

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Asn Val Tyr Val Ala Asp Leu Thr Glu Glu Val Val Tyr Thr Ser Glu 165 170 175

Met Ala Leu Lys Trp Ile Thr Lys Gly Glu Lys Ser Arg His Tyr Gly 180 185 190

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	Ile 1475					1480					Leu 1485			
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Ala His Met His Leu Lys Glu Gln Glu Thr Ile Asp Lys Leu Arg Gly Ile Val Ser Glu Lys Thr Asp Lys Leu Ser Asn Met Gln Lys Asp Leu Glu Asn Ser Asn Ala Lys Leu Gln Glu Lys Ile Gln Glu Leu Lys Ala Asn Glu His Gln Leu Ile Thr Leu Lys Lys Asp Val Asn Glu Thr Gln Lys Lys Val Ser Glu Met Glu Gln Leu Lys 1775 1780 Lys Gln Ile Lys Asp Gln Ser Leu Thr Leu Ser Lys Leu Glu Ile Glu Asn Leu Asn Leu Ala Gln Glu Leu His Glu Asn Leu Glu Glu Met Lys Ser Val Met Lys Glu Arg Asp Asn Leu Arg Arg Val Glu Glu Thr Leu Lys Leu Glu Arg Asp Gln Leu Lys Glu Ser Leu Gln Glu Thr Lys Ala Arg Asp Leu Glu Ile Gln Gln Glu Leu Lys Thr 1850 1855 Ala Arg Met Leu Ser Lys Glu His Lys Glu Thr Val Asp Lys Leu 1865 1870 Arg Glu Lys Ile Ser Glu Lys Thr Ile Gln Ile Ser Asp Ile Gln Lys Asp Leu Asp Lys Ser Lys Asp Glu Leu Gln Lys Lys Asp Arg Gln Asn His Gln Val Lys Pro Glu Lys Arg Leu Leu Ser Asp Gly

Gln Gln His Leu Met Glu Ser Leu Arg Glu Lys Cys Ser Arg Ile Lys Glu Leu Leu Lys Arg Tyr Ser Glu Met Asp Asp His Tyr Glu Cys Leu Asn Arg Leu Ser Leu Asp Leu Glu Lys Glu Ile Glu Phe His Arg Ile Met Lys Lys Leu Lys Tyr Val Leu Ser Tyr Val Thr Lys Ile Lys Glu Glu Gln His Glu Cys Ile Asn Lys Phe Glu Met Asp Phe Ile Asp Glu Val Glu Lys Gln Lys Glu Leu Leu Ile Lys Ile Gln His Leu Gln Gln Asp Cys Asp Val Pro Ser Arg Glu Leu Arg Asp Leu Lys Leu Asn Gln Asn Met Asp Leu His Ile Glu Glu Ile Leu Lys Asp Phe Ser Glu Ser Glu Phe Pro Ser Ile Lys Thr Glu Phe Gln Gln Val Leu Ser Asn Arg Lys Glu Met Thr Gln Phe Leu Glu Glu Trp Leu Asn Thr Arg Phe Asp Ile Glu Lys Leu Lys Asn Gly Ile Gln Lys Glu Asn Asp Arg Ile Cys Gln Val Asn Asn Phe Phe Asn Asn Arg Ile Ile Ala Ile Met Asn Glu Ser Thr Glu Phe Glu Glu Arg Ser Ala Thr Ile Ser Lys Glu Trp Glu Gln Asp

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Gln	Thr 2150	Leu	Lys	Thr	Ser	Leu 2155	Ala	Ser	Gly	Ala	Gln 2160	Val	Asn	Pro
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Gln	Leu 2180	Thr	Thr	Glu	Lys	Ile 2185	Arg	Glu	Leu	Glu	Asn 2190	Ser	Leu	His
Glu	Ala 2195	Lys	Glu	Ser	Ala	Met 2200	His	Lys	Glu	Ser	Lys 2205	Ile	Ile	Lys
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Pro	Leu 2330	Thr	Cys	Gly	Gly	Gly 2335	Ser	Gly	Ile	Val	Gln 2340	Asn	Thr	Lys

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